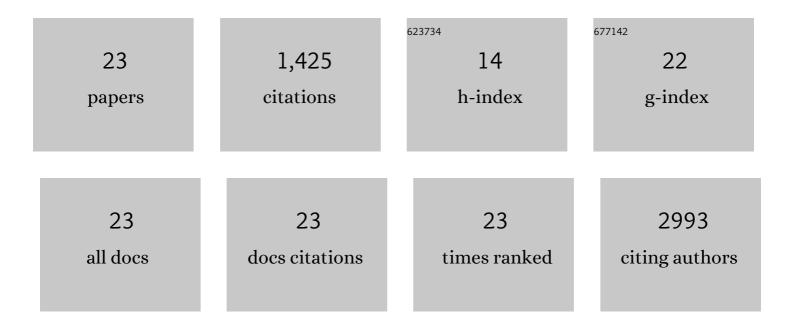
kosuke teshima

List of Publications by Year in descending order

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KOSIIKE TESHIMA

#	Article	IF	CITATIONS
1	Effects of single nucleotide polymorphism ascertainment on population structure inferences. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	6
2	Development and characterization of EST-SSR markers for <i>Pinus thunbergii</i> . Journal of Forest Research, 2021, 26, 464-467.	1.4	2
3	Lower promoter activity of the ST8SIA2 gene has been favored in evolving human collective brains. PLoS ONE, 2021, 16, e0259897.	2.5	0
4	Population genetic analysis of two species of Distylium: D. racemosum growing in East Asian evergreen broad-leaved forests and D. lepidotum endemic to the Ogasawara (Bonin) Islands. Tree Genetics and Genomes, 2019, 15, 1.	1.6	3
5	Autophagy controls reactive oxygen species homeostasis in guard cells that is essential for stomatal opening. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 19187-19192.	7.1	68
6	Field transcriptome revealed a novel relationship between nitrate transport and flowering in Japanese beech. Scientific Reports, 2019, 9, 4325.	3.3	18
7	The potential role of temperate Japanese regions as refugia for the coral Acropora hyacinthus in the face of climate change. Scientific Reports, 2019, 9, 1892.	3.3	49
8	Inferring the demographic history of Japanese cedar, Cryptomeria japonica, using amplicon sequencing. Heredity, 2019, 123, 371-383.	2.6	7
9	Potential of Genome-Wide Studies in Unrelated Plus Trees of a Coniferous Species, Cryptomeria japonica (Japanese Cedar). Frontiers in Plant Science, 2018, 9, 1322.	3.6	16
10	Effects of cyclic changes in population size on neutral genetic diversity. Ecology and Evolution, 2018, 8, 9362-9371.	1.9	4
11	Inferences of population structure and demographic history for <i>Taxodium distichum</i> , a coniferous tree in North America, based on amplicon sequencing analysis. American Journal of Botany, 2016, 103, 1937-1949.	1.7	2
12	Closely related and sympatric but not all the same: genetic variation of Indo-West Pacific Rhizophora mangroves across the Malay Peninsula. Conservation Genetics, 2015, 16, 137-150.	1.5	36
13	Inferences of evolutionary history of a widely distributed mangrove species, <i>Bruguiera gymnorrhiza</i> , in the Indoâ€West Pacific region. Ecology and Evolution, 2013, 3, 2251-2261.	1.9	35
14	The Coalescent with Selection on Copy Number Variants. Genetics, 2012, 190, 1077-1086.	2.9	20
15	Population structure and demographic history of a tropical lowland rainforest tree species <i>Shorea parvifolia</i> (Dipterocarpaceae) from Southeastern Asia. Ecology and Evolution, 2012, 2, 1663-1675.	1.9	23
16	mbs: modifying Hudson's ms software to generate samples of DNA sequences with a biallelic site under selection. BMC Bioinformatics, 2009, 10, 166.	2.6	51
17	Variations in Hd1 proteins, <i>Hd3a</i> promoters, and <i>Ehd1</i> expression levels contribute to diversity of flowering time in cultivated rice. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 4555-4560.	7.1	283
18	Natural Selection on Genes that Underlie Human Disease Susceptibility. Current Biology, 2008, 18, 883-889.	3.9	207

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#	Article	IF	CITATIONS
19	Neofunctionalization of Duplicated Genes Under the Pressure of Gene Conversion. Genetics, 2008, 178, 1385-1398.	2.9	52
20	Directional Positive Selection on an Allele of Arbitrary Dominance. Genetics, 2006, 172, 713-718.	2.9	73
21	How reliable are empirical genomic scans for selective sweeps?. Genome Research, 2006, 16, 702-712.	5.5	352
22	The Effect of Gene Conversion on the Divergence Between Duplicated Genes. Genetics, 2004, 166, 1553-1560.	2.9	106
23	The Effect of Migration During the Divergence. Theoretical Population Biology, 2002, 62, 81-95.	1.1	12